## SEQUENCE LISTING

```
<110> Diversa Corporation
      Short, Jay M.
      Kretz, Keith A.
```

<120> Recombinant Bacterial Phytases and Uses Thereof

```
<130> 81551/261654
     <140> US 09/318,528
     <141> 1999-05-25
      <150> US 09/291,931
      <151> 1999-04-13
      <150> US 09/259,214
      <151> 1999-03-01
      <150> US 08/910,798
      <151> 1997-08-13
      <160> 4
      <170> FastSEQ for Windows Version 3.0
     <210> 1
      <211> 1323
      <212> DNA
      <213> Escherichia coli
      <220>
      <221> CDS
      <222> (1)...(1320)
      <400> 1
atg aaa gcg atc tta atc cca ttt tta tct ctt ctg att ccg tta acc
                                                                        48
Met Lys Ala Ile Leu Ile Pro Phe Leu Ser Leu Leu Ile Pro Leu Thr
                 5
                                      10
                                                          15
ccg caa tct gca ttc gct cag agt gag ccg gag ctg aag ctg gaa agt
                                                                        96
Pro Gln Ser Ala Phe Ala Gln Ser Glu Pro Glu Leu Lys Leu Glu Ser
             20
                                  25
gtg gtg att gtc agt cgt cat ggt gtg cgt gct cca acc aag gcc acg
```

Val Val Ile Val Ser Arg His Gly Val Arg Ala Pro Thr Lys Ala Thr

40

35

144

caa Gln	ctg Leu 50	atg Met	cag Gln	gat Asp	gtc Val	acc Thr 55	cca Pro	gac Asp	gca Ala	tgg Trp	cca Pro 60	acc Thr	tgg Trp	ccg Pro	gta Val	192
aaa Lys 65	ctg Leu	ggt Gly	tgg Trp	ctg Leu	aca Thr 70	ccg Pro	cgn Xaa	ggt Gly	ggt Gly	gag Glu 75	cta Leu	atc Ile	gcc Ala	tat Tyr	ctc Leu 80	240
gga Gly	cat His	tac Tyr	caa Gln	cgc Arg 85	cag Gln	cgt Arg	ctg Leu	gta Val	gcc Ala 90	gac Asp	gga Gly	ttg Leu	ctg Leu	gcg Ala 95	aaa Lys	288
aag Lys	ggc Gly	tgc Cys	ccg Pro 100	cag Gln	tct Ser	ggt Gly	cag Gln	gtc Val 105	gcg Ala	att Ile	att Ile	gct Ala	gat Asp 110	gtc Val	gac Asp	336
gag Glu	cgt Arg	acc Thr 115	cgt Arg	aaa Lys	aca Thr	ggc Gly	gaa Glu 120	gcc Ala	ttc Phe	gcc Ala	gcc Ala	ggg Gly 125	ctg Leu	gca Ala	cct Pro	384
gac Asp	tgt Cys 130	gca Ala	ata Ile	acc Thr	gta Val	cat His 135	acc Thr	cag Gln	gca Ala	gat Asp	acg Thr 140	tcc Ser	agt Ser	ccc Pro	gat Asp	432
ccg Pro 145	Leu	ttt Phe	aat Asn	cct Pro	cta Leu 150	aaa Lys	act Thr	ggc Gly	gtt Val	tgc Cys 155	caa Gln	ctg Leu	gat Asp	aac Asn	gcg Ala 160	480
aac Asn	gtg Val	act Thr	gac Asp	gcg Ala 165	atc Ile	ctc Leu	agc Ser	agg Arg	gca Ala 170	gga Gly	ggg Gly	tca Ser	att Ile	gct Ala 175	gac Asp	528
ttt Phe	acc Thr	ggg	cat His 180	cgg Arg	caa Gln	acg Thr	gcg Ala	ttt Phe 185	cgc Arg	gaa Glu	ctg Leu	gaa Glu	cgg Arg 190	gtg Val	ctt Leu	576
aat Asn	ttt Phe	ccg Pro 195	caa Gln	tca Ser	aac Asn	ttg Leu	tgc Cys 200	ctt Leu	aaa Lys	cgt Arg	gag Glu	aaa Lys 205	cag Gln	gac Asp	gaa Glu	624
agc Ser	tgt Cys 210	tca Ser	tta Leu	acg Thr	cag Gln	gca Ala 215	tta Leu	cca Pro	tcg Ser	gaa Glu	ctc Leu 220	aag Lys	gtg Val	agc Ser	gcc Ala	672
gac Asp 225	aat Asn	gtc Val	tca Ser	tta Leu	acc Thr 230	ggt Gly	gcg Ala	gta Val	agc Ser	ctc Leu 235	gca Ala	tca Ser	atg Met	ctg Leu	acg Thr 240	720
gag Glu	ata Ile	ttt Phe	ctc Leu	ctg Leu 245	caa Gln	caa Gln	gca Ala	cag Gln	gga Gly 250	atg Met	ccg Pro	gag Glu	ccg Pro	ggg Gly 255	tgg Trp	768

		atc Ile														816
		caa Gln 275														864
		acc Thr														912
		caa Gln														960
		gcc Ala														1008
		aac Asn														1056
		ctg Leu 355														1104
		cag Gln														1152
		ccg Pro														1200
ctg Leu	gca Ala	gga Gly	tgt Cys	gaa Glu 405	gag Glu	cga Arg	aat Asn	gcg Ala	cag Gln 410	ggc Gly	atg Met	tgt Cys	tcg Ser	ttg Leu 415	gca Ala	1248
ggt Gly	ttt Phe	acg Thr	caa Gln 420	atc Ile	gtg Val	aat Asn	gaa Glu	gca Ala 425	cgc Arg	ata Ile	ccg Pro	gcg Ala	tgc Cys 430	agt Ser	ttg Leu	1296
		cat His 435						taa								1323

```
<210> 2
<211> 440
<212> PRT
<213> Escherichia coli
```

325

355

<400> 2 Met Lys Ala Ile Leu Ile Pro Phe Leu Ser Leu Leu Ile Pro Leu Thr Pro Gln Ser Ala Phe Ala Gln Ser Glu Pro Glu Leu Lys Leu Glu Ser Val Val Ile Val Ser Arg His Gly Val Arg Ala Pro Thr Lys Ala Thr 40 Gln Leu Met Gln Asp Val Thr Pro Asp Ala Trp Pro Thr Trp Pro Val Lys Leu Gly Trp Leu Thr Pro Xaa Gly Gly Glu Leu Ile Ala Tyr Leu 75 Gly His Tyr Gln Arg Gln Arg Leu Val Ala Asp Gly Leu Leu Ala Lys 85 90 Lys Gly Cys Pro Gln Ser Gly Gln Val Ala Ile Ile Ala Asp Val Asp 105 110 Glu Arg Thr Arg Lys Thr Gly Glu Ala Phe Ala Ala Gly Leu Ala Pro 120 Asp Cys Ala Ile Thr Val His Thr Gln Ala Asp Thr Ser Ser Pro Asp 135 Pro Leu Phe Asn Pro Leu Lys Thr Gly Val Cys Gln Leu Asp Asn Ala 150 155 Asn Val Thr Asp Ala Ile Leu Ser Arg Ala Gly Gly Ser Ile Ala Asp 165 170 Phe Thr Gly His Arg Gln Thr Ala Phe Arg Glu Leu Glu Arg Val Leu 180 185 Asn Phe Pro Gln Ser Asn Leu Cys Leu Lys Arg Glu Lys Gln Asp Glu 200 Ser Cys Ser Leu Thr Gln Ala Leu Pro Ser Glu Leu Lys Val Ser Ala 215 220 Asp Asn Val Ser Leu Thr Gly Ala Val Ser Leu Ala Ser Met Leu Thr 230 235 Glu Ile Phe Leu Leu Gln Gln Ala Gln Gly Met Pro Glu Pro Gly Trp 245 250 Gly Arg Ile Thr Asp Ser His Gln Trp Asn Thr Leu Leu Ser Leu His 260 265 Asn Ala Gln Phe Tyr Leu Leu Gln Arg Thr Pro Glu Val Ala Arg Ser 280 Arg Ala Thr Pro Leu Leu Asp Leu Ile Met Ala Ala Leu Thr Pro His 295 300 Pro Pro Gln Lys Gln Ala Tyr Gly Val Thr Leu Pro Thr Ser Val Leu 310 315 Phe Ile Ala Gly His Asp Thr Asn Leu Ala Asn Leu Gly Gly Ala Leu

365

330

Glu Leu Asn Trp Thr Leu Pro Gly Gln Pro Asp Asn Thr Pro Pro Gly 340 345 350 Gly Glu Leu Val Phe Glu Arg Trp Arg Leu Ser Asp Asn Ser Gln

360

	Trp	Ile 370	Gln	Val	Ser	Leu	Val 375	Phe	Gln	Thr	Leu	Gln 380	Gln	Met	Arg	Asp	
	Lys 385	Thr	Pro	Leu	Ser	Leu 390	Asn	Thr	Pro	Pro	Gly 395	Glu	Val	Lys	Leu	Thr 400	
	Leu	Ala	Gly	Суз	Glu 405	Glu	Arg	Asn	Ala	Gln 410		Met	Cys	Ser	Leu 415		
	Gly	Phe	Thr	Gln 420	Ile	Val	Asn	Glu	Ala 425		Ile	Pro	Ala	Cys 430	Ser	Leu	
	Arg	Ser	His 435	His	His	His	His	His 440									
		_		_													
	<210> 3																
	<211> 49																
	<212> DNA																
	<213> Artificial Sequence																
	<220>																
		<2	223>	Olig	gonu	cleot	ide										
		< 1	100>	3													
	atti				raaa	ga at	++==	aataa	2 22/	rcast	+	22+	2001	- +-			4.0
	9000	Jocyc	146 (	Juan	gag	ga a	Luca	aacy	ı aay	jcya	-000	aatt	Jula	- (			49
		<2	210>	4													
			211>														
			212>														
					ific	ial S	Seaue	ence									
							- 1										
		<2	220>														
		<2	223>	Oli	gonu	cleot	cide										
		<4	100>	4													
	gttt				acaaa	ac to	caco	accad	r tat	_							33
			-						-								

33